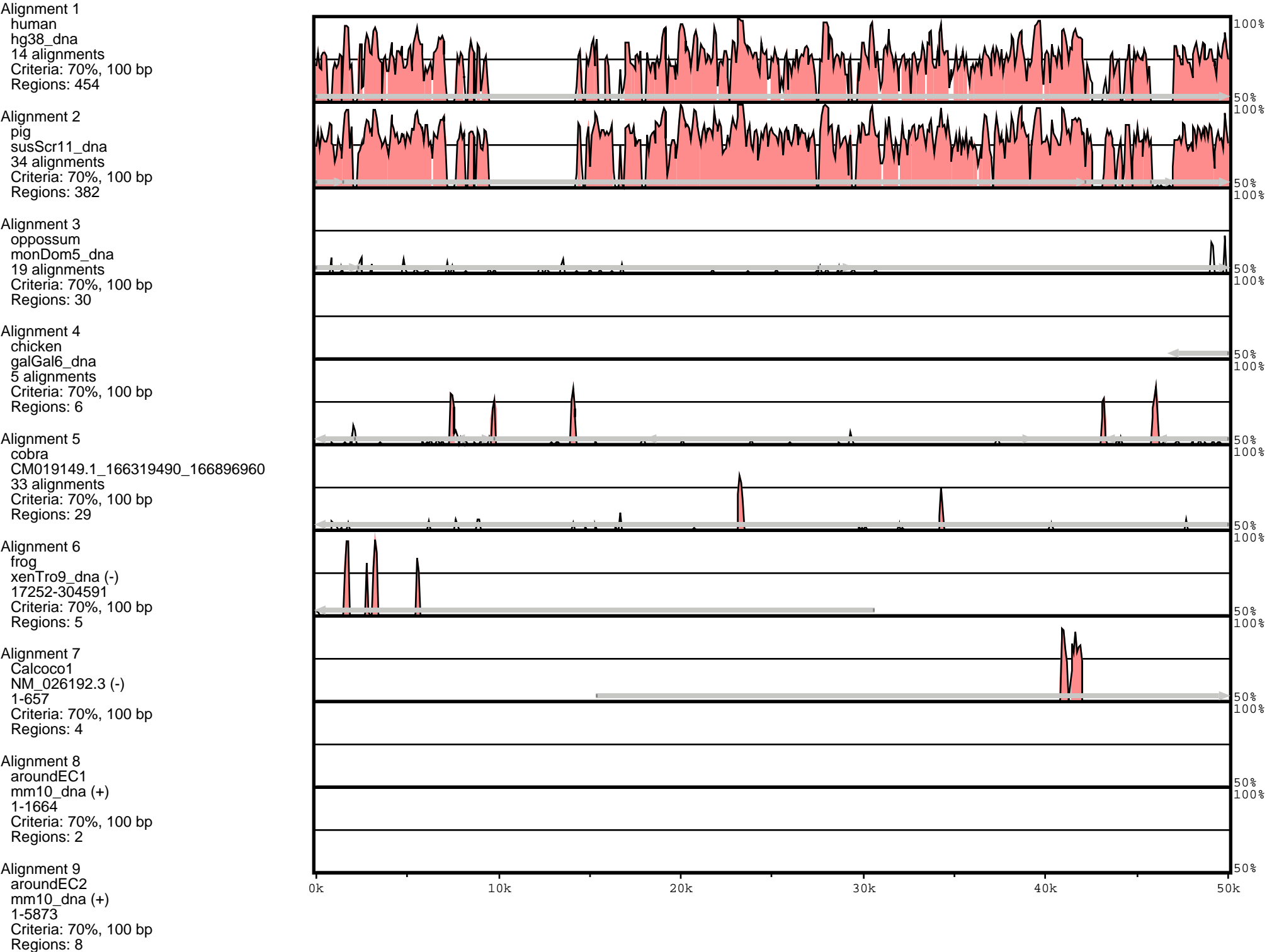
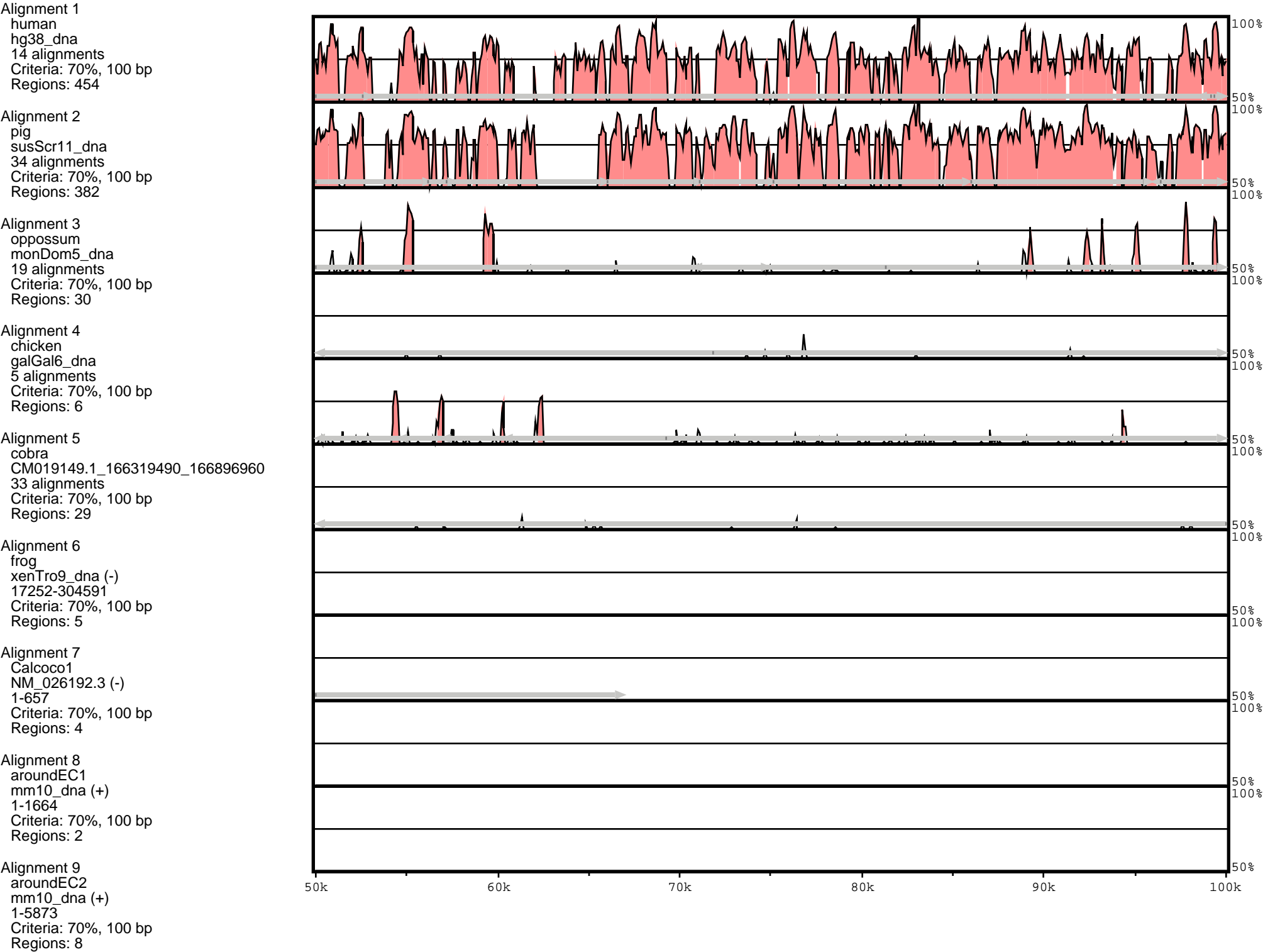


sheep oviAri4\_dna:1-250530



sheep oviAri4\_dna:1-250530



Alignment 1  
human  
hg38\_dna  
14 alignments  
Criteria: 70%, 100 bp  
Regions: 454

Alignment 2  
pig  
susScr11\_dna  
34 alignments  
Criteria: 70%, 100 bp  
Regions: 382

Alignment 3  
opposum  
monDom5\_dna  
19 alignments  
Criteria: 70%, 100 bp  
Regions: 30

Alignment 4  
chicken  
galGal6\_dna  
5 alignments  
Criteria: 70%, 100 bp  
Regions: 6

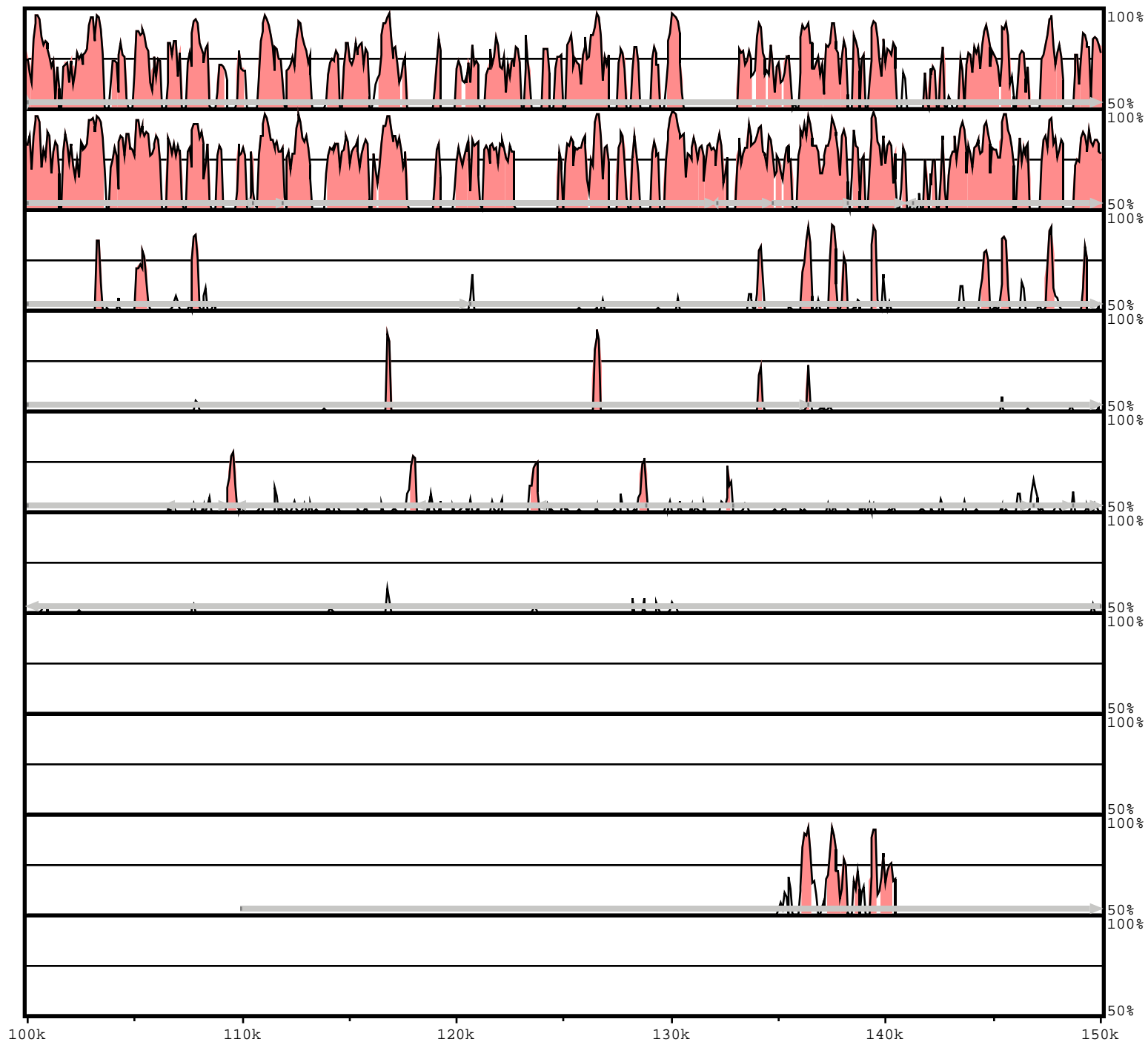
Alignment 5  
cobra  
CM019149.1\_166319490\_166896960  
33 alignments  
Criteria: 70%, 100 bp  
Regions: 29

Alignment 6  
frog  
xenTro9\_dna (-)  
17252-304591  
Criteria: 70%, 100 bp  
Regions: 5

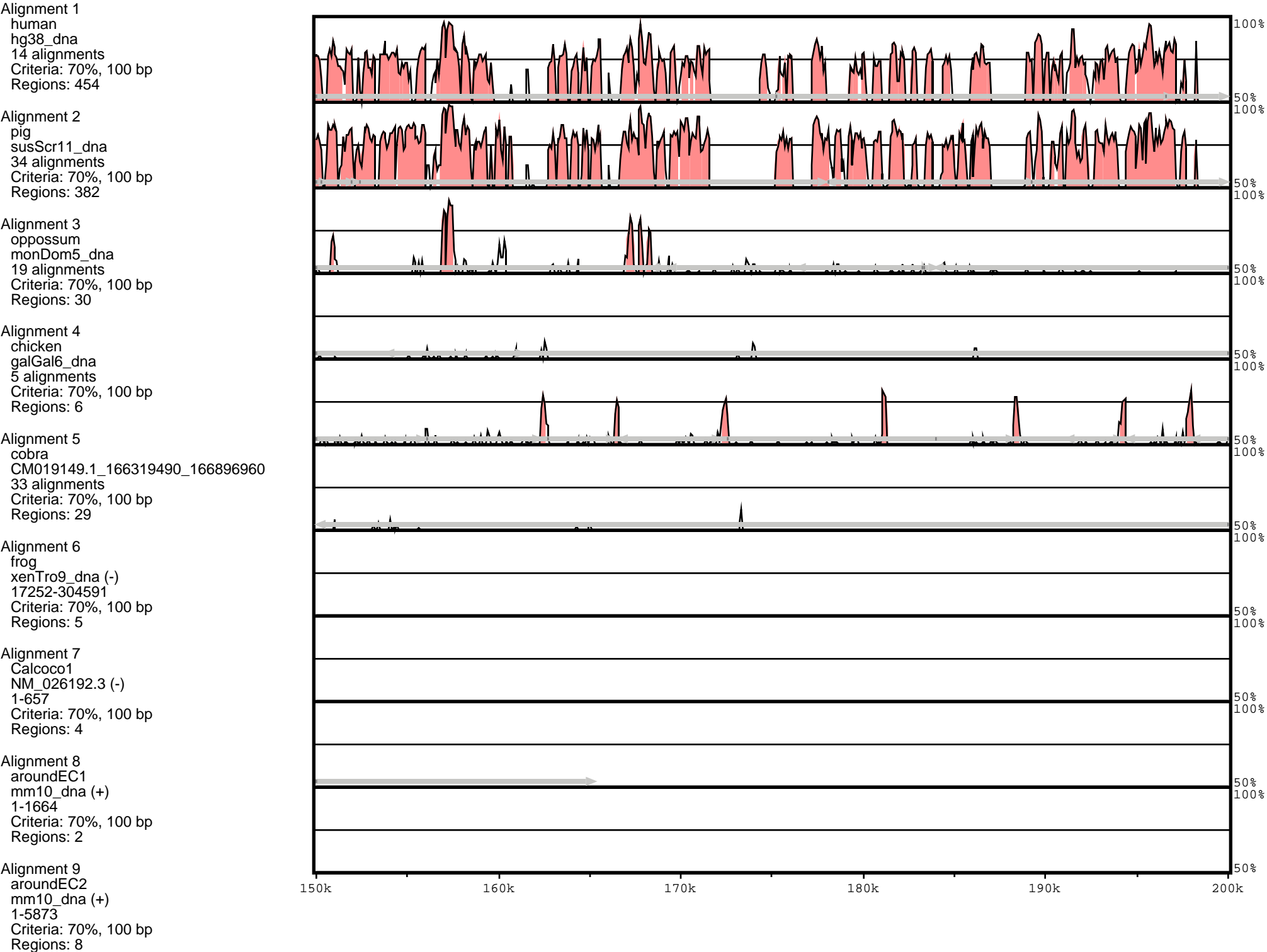
Alignment 7  
Calcoco1  
NM\_026192.3 (-)  
1-657  
Criteria: 70%, 100 bp  
Regions: 4

Alignment 8  
aroundEC1  
mm10\_dna (+)  
1-1664  
Criteria: 70%, 100 bp  
Regions: 2

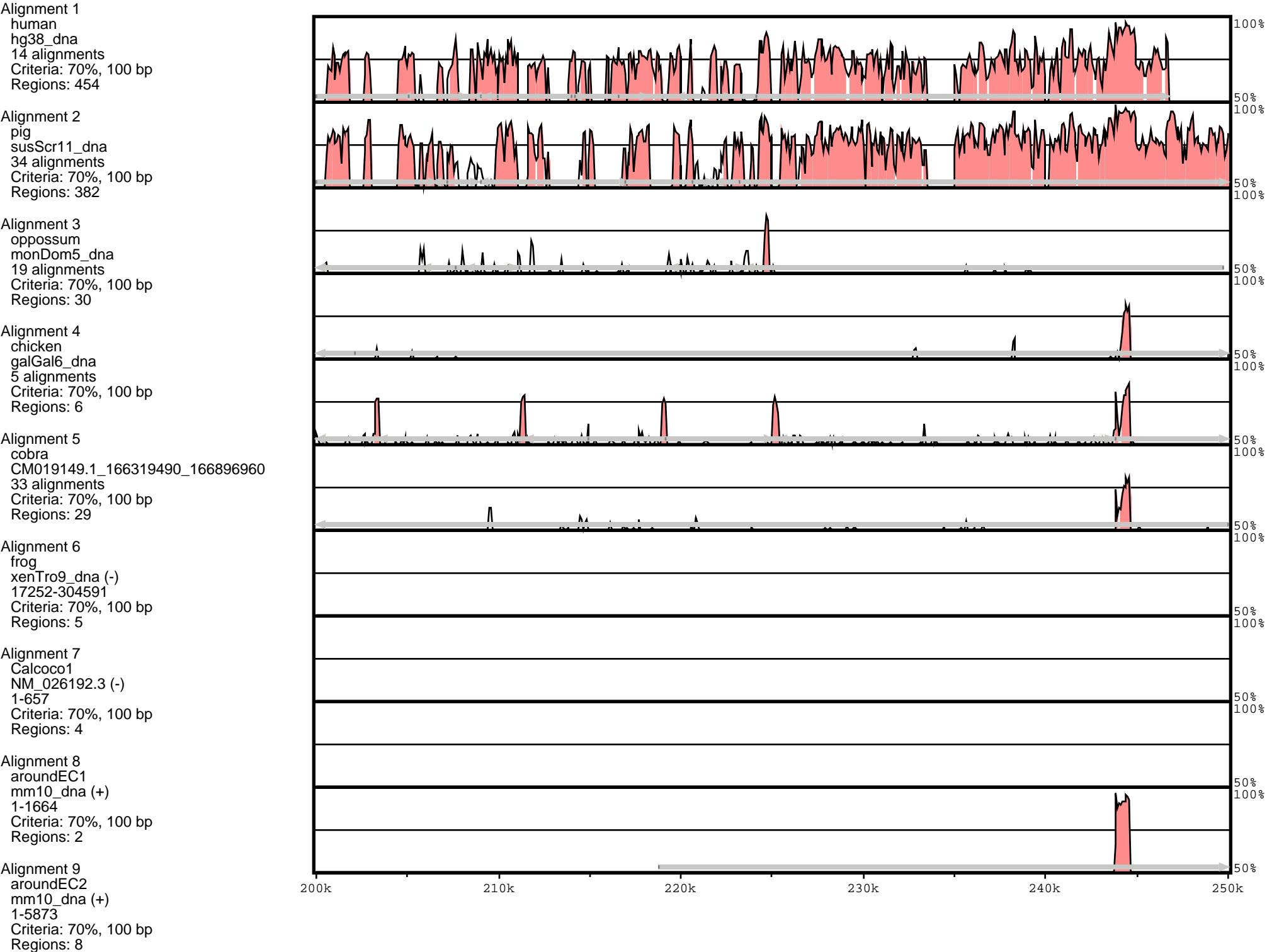
Alignment 9  
aroundEC2  
mm10\_dna (+)  
1-5873  
Criteria: 70%, 100 bp  
Regions: 8



sheep oviAri4\_dna:1-250530



sheep oviAri4\_dna:1-250530



Alignment 1  
human  
hg38\_dna  
14 alignments  
Criteria: 70%, 100 bp  
Regions: 454

Alignment 3  
opossum  
monDom5\_dna  
19 alignments  
Criteria: 70%, 100 bp  
Regions: 30

Alignment 4  
chicken  
galGal6\_dna  
5 alignments  
Criteria: 70%, 100 bp  
Regions: 6

Alignment 5  
cobra  
CM019149.1\_166319490\_166896960  
33 alignments  
Criteria: 70%, 100 bp  
Regions: 29

Alignment 6  
frog  
xenTro9\_dna (-)  
17252-304591  
Criteria: 70%, 100 bp  
Regions: 5

Alignment 7  
Calcoco1  
NM\_026192.3 (-)  
1-657  
Criteria: 70%, 100 bp  
Regions: 4

Alignment 8  
around EC1  
mm10\_dna (+)  
1-1664  
Criteria: 70%, 100 bp  
Regions: 2

Alignment 9  
around EC2  
mm10\_dna (+)  
1-5873  
Criteria: 70%, 100 bp  
Regions: 8

